

CURRICULUM VITAE

NAME THAN VAN THAI D.V.M, Ph.D.	CURRENT POSITION Postdoctoral Fellow Chung-Ang University Department of Microbiology, College of Medicine 221 Huksuk-Dong, Dongjak-Ku Seoul 156-756 Korea E-mail: thanthaivnn@yahoo.com Phone: +82 10-9109-2268
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A. EDUCATION

INSTITUTION	DEGREE	YEAR	FIELD OF STUDY
Hanoi Agriculture University, Vietnam	D.V.M.	2006	Veterinary Medicine
Chung-Ang university, South Korea	M.Sc.	2011	Virology
Chung-Ang university, South Korea	Ph.D.	2014	Virology

B. EMPLOYMENTS

ORGANIZATION	POSITION	YEAR
Hanoi Agriculture University, Vietnam	D.V.M. Student	2001-2006
National Centre for Veterinary Diagnostics, Vietnam	Staff Scientist	2006-2009
Chung-Ang University, South Korea	M.Sc. Student	2009-2011
Chung-Ang University, South Korea	Ph.D. Student	2011-2014
Chung-Ang University, South Korea	Postdoctoral Fellow	2014-Present

C. HONORS

2009-2011: Chung-Ang Young Scientist Scholarship for M.Sc. Student, South Korea

2011-2014: Chung-Ang Young Scientist Scholarship for Ph.D. Student, South Korea

D. PEER-REVIEWED ACTIVITIES FOR PEER-REVIEWED JOURNALS

Journal of Medical Virology

BMC Infectious Diseases

PLOS ONE

Journal of Microbiology and Biotechnology

Epidemiology and Infection

Archives of Virology

E. RESEARCH INTERESTS

Novel Virus Discovery

Viral Evolution, Epidemiology, Emerging Infectious Diseases

Gene Expression in Baculovirus System, Protein Purification, Virus-like Particle

Immunology

F. PUBLICATIONS

1. **Than VT**, Jeong S, Kim W. 2014. Characterization of rotaTeq[®] vaccine-derived rotaviruses in South Korean infants with rotavirus gastroenteritis. J Med Virol. PMID: 24817171 [Epub ahead of print].
2. Jeong S, **Than VT**, Lim I, Kim W. 2014. Whole-Genome analysis of a rare human Korean G3P[9] rotavirus strain suggests a complex evolutionary origin potentially involving reassortment events between feline and bovine rotaviruses. PLoS One. 9(5):e97127.
3. Thoi TC, **Than VT**, Kim W. 2014. Whole genomic characterization of a Korean human parechovirus type 1 (HPeV1) identifies recombination events. J Med Virol. PMID: 24634137 [Epub ahead of print].
4. **Than VT**, Baek IH, Kim W. 2013. Whole genomic analysis reveals the co-evolutionary phylodynamics of Korean G9P[8] human rotavirus strains. Arch Virol. 158(8):1795-803.
5. **Than VT**, Kang H, Lim I, Kim W. 2013. Molecular characterization of serotype G9 rotaviruses circulating in South Korea between 2005 and 2010. J Med Virol. 85(1):171-8.
6. **Than VT**, Park JH, Chung IS, Kim JB, Kim W. 2013. Whole-genome sequence analysis of a Korean G11P[25] rotavirus strain identifies several porcine-human reassortant events. Arch Virol. 158(11):2385-93.
7. **Than VT**, Kim W. 2013. Prevalence of rotavirus genotypes in South Korea in 1989-2009: implications for a nationwide rotavirus vaccine program. Korean J Pediatr. 56(11):465-473.

8. Shim JO, **Thai VT**, Ryoo E, Lim I, Yoon Y, Kim K, Chung SI, Kim W. 2013. Distribution of rotavirus G and P genotypes approximately two years following the introduction of rotavirus vaccines in South Korea. *J Med Virol.* 85(7):1307-12.
9. Baek IH, **Than VT**, Kim H, Lim I, Kim W. 2013. Full genomic characterization of a group C rotavirus isolated from a child in South Korea. *J Med Virol.* 85(8):1478-84.
10. **Than VT**, Baek IH, Lee HY, Kim JB, Shon DH, Chung IS, Kim W. 2012. Expression of recombinant rotavirus proteins harboring antigenic epitopes of the hepatitis A virus polypeptide in insect cells. *Biomol Ther.* 20(3):320-5.
11. Shin JW, Le VP, **Than VT**, Lim I, Yoon Y, Kim K, Chung SI, Myung SC, Kim W. 2012. Genomic characterization of a cell-culture-adapted Korean human G9P[8] rotavirus, CAU05-202. *Arch Virol.* 157(4):753-9.
12. **Than VT**, Le VP, Lim I, Kim W. 2011. Complete genomic characterization of cell culture adapted human G12P[6] rotaviruses isolated from South Korea. *Virus Genes.* 42(3):317-22.